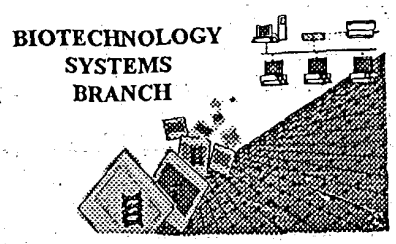

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/462,480A
Source: IFW16
Date Processed by STIC: 11/23/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) **INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
2) **TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , **EFS Submission User Manual** - ePAVE).
2. **U.S. Postal Service:** Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. **Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):** U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04

Raw Sequence Listing Error Summary

ERROR DETECTED **SUGGESTED CORRECTION** **SERIAL NUMBER:** 09/462,480A

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

1	Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor **after** creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2	Invalid Line Length	The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
3	Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
4	Non-ASCII	The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. **Please ensure your subsequent submission is saved in ASCII text.**
5	Variable Length	Sequence(s)____contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6	PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
7	Skipped Sequences (OLD RULES)	Sequence(s) _____ missing. If intentional, please insert the following lines for **each** skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to **include** the skipped sequences.
8	Skipped Sequences (NEW RULES)	Sequence(s) _____ missing. **If intentional**, please insert the following lines for **each** skipped sequence. <210> sequence id number <400> sequence id number 000
9	Use of n's or Xaa's (NEW RULES)	Use of **n**'s and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of **n** or **Xaa**, and which residue **n** or **Xaa** represents.
10	Invalid <213> Response	Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence. Do not combine responses.
11	Use of <220>	Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
12	PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
13	Misuse of n/Xaa	"n" can **only** represent a single nucleotide; "Xaa" can **only** represent a single amino acid

IFW16

RAW SEQUENCE LISTING DATE: 11/23/2004
PATENT APPLICATION: **US/09/462,480A** TIME: 14:08:31

Input Set : **A:\066001650XPCT.txt**
Output Set: **N:\CRF4\11222004\I462480A.raw**

<110> APPLICANT: GICQUEL, BRIGITTE
BERTHET, FRANCIOS-XAVIER
ANDERSEN, PETER
RASMUSSEN, PETER BIRK
<120> TITLE OF INVENTION: POLYNUCLEOTIDE FUNCTIONALLY CODING FOR THE LHP PROTEIN FROM MYCOBACTERIUM TUBERCULOSIS, ITS BIOLOGICALLY ACTIVE DERIVATIVE FRAGMENTS, AS WELL AS METHODS USING THE SAME
<130> FILE REFERENCE: 0660-0165-0XPCT
<140> CURRENT APPLICATION NUMBER: 09/462,480A
<141> CURRENT FILING DATE: 2000-03-06
<150> PRIOR APPLICATION NUMBER: PCT/IB98/01091
<151> PRIOR FILING DATE: 1998-07-16
<150> PRIOR APPLICATION NUMBER: 60/052,631
<151> PRIOR FILING DATE: 1997-07-16
<160> NUMBER OF SEQ ID NOS: 34
<170> SOFTWARE: PatentIn version 3.3
<210> SEQ ID NO: 1
<211> LENGTH: 1277
<212> TYPE: DNA
<213> ORGANISM: Mycobacterium tuberculosis
<400> SEQUENCE: 1

pr 4-6

Does Not Comply
Corrected Diskette Needed

ctgcagcagg tgacgtcgtt gttcagccag gtgggcggca ccggcggcgg caacccagcc 60
gacgaggaag ccgcgcagat gggcctgctc ggcaccagtc cgctgtcgaa ccatccgctg 120
gctggtggat caggccccag cgcgggcgcg ggcctgctgc gcgcggagtc gctacctggc 180
gcaggtgggt cgttgacccg cacgccgctg atgtctcagc tgatcgaaaa gccggttgcc 240
ccctcggtga tgccggcggc tgttgccgga tcgtcggtga cgggtggcgc cgctccggtg 300
ggtccgggag cgatgggcca gggttcgcaa tccggcggct ccaccagccc gggtctggtc 360
gcgccggcac cgctcgcgca ggagcgtgaa gaagacgacg aggacgactg ggacgaagag 420
gacgactggt gagctcccgt aatgacaaca gacttcccgg ccacccgggc cggaagactt 480
gccaacattt tggcgaggaa ggtaaagaga gaaagtagtc cagcatggca gagatgaaga 540
ccgatgccgc taccctcggg caggaggcag gtaatttcga gcggatctcc ggcgacctga 600
aaacccagat cgaccaggtg gagtcgacgg caggttcgtt gcagggccag tggcgcggcg 660
cggcggggac ggccgcccag gccgcggtgg tgcgcttcca agaagcagcc aataagcaga 720
agcaggaact cgacgagatc tcgacgaata ttcgtcaggc cggcgtccaa tactcgaggg 780
ccgacgagga gcagcagcag gcgctgtcct cgcaaatggg cttctgaccc gctaatacga 840
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ggcaagcgca atccagggaa atgtcacgtc cattcattcc ctccttgacg aggggaagca 960
gtccctgacc aagctcgcag cggcctgggg cggtagcggt tcggaggcgt accagggtgt 1020
ccagcaaaaa tgggacgcca cggctaccga gctgaacaac gcgctgcaga acctggcgcg 1080
gacgatcagc gaagccggtc aggcaatggc ttcgaccgaa ggcaacgtca ctgggatgtt 1140
cgcatagggc aacgccgagt tcgcgtagaa tagcgaaaca cgggatcggg cgagttcgac 1200
cttccgtcgg tctcgccctt tctcgtgttt atacgtttga gcgcactctg agaggttgtc 1260
atggcggccg actacga 1277

RAW SEQUENCE LISTING DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A TIME: 14:08:31

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

<210> SEQ ID NO: 2
<211> LENGTH: 524
<212> TYPE: DNA
<213> ORGANISM: Mycobacterium tuberculosis
<400> SEQUENCE: 2
ctgcagcagg tgacgtcgtt gttcagccag gtgggcggca ccggcggcgg caacccagcc 60
gacgaggaag ccgcgcagat gggcctgctc ggcaccagtc cgctgtcgaa ccatccgctg 120
gctggtggat caggccccag cgcgggcgcg ggcctgctgc gcgcggagtc gctacctggc 180
gcaggtgggt cgttgacccg cacgccgctg atgtctcagc tgatcgaaaa gccggttgcc 240
ccctcggtga tgccggcggc tgttgccgga tcgtcggtga cgggtggcgc cgctccggtg 300
ggtccgggag cgatgggcca gggttcgcaa tccggcggct ccaccagccc gggtctggtc 360
gcgccggcac cgctcgcgca ggagcgtgaa gaagacgacg aggacgactg ggacgaagag 420
gacgactggt gagctcccgt aatgacaaca gacttcccgg ccacccgggc cggaagactt 480
gccaacattt tggcgaggaa ggtaaagaga gaaagtagtc cagc 524
<210> SEQ ID NO: 3
<211> LENGTH: 481
<212> TYPE: DNA
<213> ORGANISM: Mycobacterium tuberculosis
<400> SEQUENCE: 3
ctgcagcagg tgacgtcgtt gttcagccag gtgggcggca ccggcggcgg caacccagcc 60
gacgaggaag ccgcgcagat gggcctgctc ggcaccagtc cgctgtcgaa ccatccgctg 120
gctggtggat caggccccag cgcgggcgcg ggcctgctgc gcgcggagtc gctacctggc 180
gcaggtgggt cgttgacccg cacgccgctg atgtctcagc tgatcgaaaa gccggttgcc 240
ccctcggtga tgccggcggc tgttgccgga tcgtcggtga cgggtggcgc cgctccggtg 300
ggtccgggag cgatgggcca gggttcgcaa tccggcggct ccaccagccc gggtctggtc 360
gcgccggcac cgctcgcgca ggagcgtgaa gaagacgacg aggacgactg ggacgaagag 420
gacgactggt gagctcccgt aatgacaaca gacttcccgg ccacccgggc cggaagactt 480
g 481
<210> SEQ ID NO: 4
<211> LENGTH: 302
<212> TYPE: DNA
<213> ORGANISM: Mycobacterium tuberculosis
<400> SEQUENCE: 4
atggcagaga tgaagaccga tgccgctacc ctcgggcagg aggcaggtaa tttcgagcgg 60
atctccggcg acctgaaaac ccagatcgac caggtggagt cgacggcagg ttcgttgcag 120
ggccagtggc gcggcgcggc ggggacggcc gcccaggccg cggtggtgcg cttccaagaa 180
gcagccaata agcagaagca ggaactcgac gagatctcga cgaatattcg tcaggccggc 240
gtccaatact cgagggccga cgaggagcag cagcaggcgc tgtcctcgca aatgggcttc 300
tg 302
<210> SEQ ID NO: 5
<211> LENGTH: 100
<212> TYPE: PRT
<213> ORGANISM: Mycobacterium tuberculosis
<400> SEQUENCE: 5
Met Ala Glu Met Lys Thr Asp Ala Ala Thr Leu Gly Gln Glu Ala Gly
1 5 10 15
Asn Phe Glu Arg Ile Ser Gly Asp Leu Lys Thr Gln Ile Asp Gln Val
20 25 30
Glu Ser Thr Ala Gly Ser Leu Gln Gly Gln Trp Arg Gly Ala Ala Gly

RAW SEQUENCE LISTING DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A TIME: 14:08:31

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

```
163         35                  40                  45
166 Thr Ala Ala Gln Ala Ala Val Val Arg Phe Gln Glu Ala Ala Asn Lys
167     50                  55                  60
170 Gln Lys Gln Glu Leu Asp Glu Ile Ser Thr Asn Ile Arg Gln Ala Gly
171 65                  70                  75                  80
174 Val Gln Tyr Ser Arg Ala Asp Glu Glu Gln Gln Gln Ala Leu Ser Ser
175                 85                  90                  95
178 Gln Met Gly Phe
179             100
182 <210> SEQ ID NO: 6
183 <211> LENGTH: 49
184 <212> TYPE: PRT
185 <213> ORGANISM: Mycobacterium tuberculosis
187 <400> SEQUENCE: 6
189 Met Ala Glu Met Lys Thr Asp Ala Ala Thr Leu Gly Gln Glu Ala Gly
190 1               5                   10                  15
193 Asn Phe Glu Arg Ile Ser Gly Asp Leu Lys Thr Gln Ile Asp Gln Val
194             20                  25                  30
197 Glu Ser Thr Ala Gly Ser Leu Gln Gly Gln Trp Arg Gly Ala Ala Gly
198         35                  40                  45
201 Thr
205 <210> SEQ ID NO: 7
206 <211> LENGTH: 42
207 <212> TYPE: PRT
208 <213> ORGANISM: Mycobacterium tuberculosis
210 <400> SEQUENCE: 7
212 Gln Glu Ala Ala Asn Lys Gln Lys Gln Glu Leu Asp Gly Ile Ser Thr
213 1               5                   10                  15
216 Asn Ile Arg Gln Ala Gly Val Gln Tyr Ser Arg Ala Asp Glu Glu Gln
217             20                  25                  30
220 Gln Gln Ala Leu Ser Ser Gln Met Gly Phe
221         35                  40
224 <210> SEQ ID NO: 8
225 <211> LENGTH: 21
226 <212> TYPE: PRT
227 <213> ORGANISM: Mycobacterium tuberculosis
229 <400> SEQUENCE: 8
231 Gln Glu Ala Gly Asn Phe Glu Arg Ile Ser Gly Asp Leu Lys Tyr Thr
232 1               5                   10                  15
235 Gln Ile Asp Gln Val
236             20
239 <210> SEQ ID NO: 9
240 <211> LENGTH: 16
241 <212> TYPE: PRT
242 <213> ORGANISM: Mycobacterium tuberculosis
244 <400> SEQUENCE: 9
246 Gly Asp Leu Lys Thr Gln Ile Asp Gln Val Glu Ser Thr Ala Gly Ser
247 1               5                   10                  15
250 <210> SEQ ID NO: 10
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RAW SEQUENCE LISTING DATE: 11/23/2004
PATENT APPLICATION: **US/09/462,480A** TIME: 14:08:31

Input Set : **A:\066001650XPCT.txt**
Output Set: **N:\CRF4\11222004\I462480A.raw**

```
251 <211> LENGTH: 16
252 <212> TYPE: PRT
253 <213> ORGANISM: Mycobacterium tuberculosis
255 <400> SEQUENCE: 10
257 Gly Ser Leu Gln Gly Gln Trp Arg Gly Ala Ala Gly Thr Ala Ala Gln
258 1               5                   10                  15
261 <210> SEQ ID NO: 11
262 <211> LENGTH: 16
263 <212> TYPE: PRT
264 <213> ORGANISM: Mycobacterium tuberculosis
266 <400> SEQUENCE: 11
268 Gln Glu Ala Ala Asn Lys Gln Lys Gln Glu Leu Asp Glu Ile Ser Thr
269 1               5                   10                  15
272 <210> SEQ ID NO: 12
273 <211> LENGTH: 28
274 <212> TYPE: PRT
275 <213> ORGANISM: Mycobacterium tuberculosis
277 <400> SEQUENCE: 12
279 Ser Thr Asn Ile Arg Gln Ala Gly Val Gln Tyr Ser Arg Ala Asp Glu
280 1               5                   10                  15
283 Glu Gln Gln Gln Ala Leu Ser Ser Gln Met Gly Phe
284             20                  25
287 <210> SEQ ID NO: 13
288 <211> LENGTH: 16
289 <212> TYPE: PRT
290 <213> ORGANISM: Mycobacterium tuberculosis
292 <400> SEQUENCE: 13
294 Arg Ala Asp Glu Glu Gln Gln Gln Ala Leu Ser Ser Gln Met Gly Phe
295 1               5                   10                  15
298 <210> SEQ ID NO: 14
299 <211> LENGTH: 21
300 <212> TYPE: DNA
C--> 301 <213> ORGANISM: Artificial/Unknown
304 <220> FEATURE:
305 <221> NAME/KEY: misc_feature
306 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
308 <400> SEQUENCE: 14
309 ctgcagcagg tgacgtcgtt g                                          21
312 <210> SEQ ID NO: 15
313 <211> LENGTH: 23
314 <212> TYPE: DNA
C--> 315 <213> ORGANISM: Artificial/Unknown
318 <220> FEATURE:
319 <221> NAME/KEY: misc_feature
320 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
322 <400> SEQUENCE: 15
323 ccgggtggcc gggaagtctg tgt                                        23
326 <210> SEQ ID NO: 16
327 <211> LENGTH: 23
```

invalid - see item 10 on Error Summary Sheet

same error

RAW SEQUENCE LISTING DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A TIME: 14:08:31

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

```
     328 <212> TYPE: DNA
C--> 329 <213> ORGANISM: Artificial/Unknown
     332 <220> FEATURE:
     333 <221> NAME/KEY: misc_feature
     334 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
     336 <400> SEQUENCE: 16
     337 actactttct ctttctacct tcc                                          23
     340 <210> SEQ ID NO: 17
     341 <211> LENGTH: 39
     342 <212> TYPE: DNA
C--> 343 <213> ORGANISM: Artificial/Unknown
     346 <220> FEATURE:
     347 <221> NAME/KEY: misc_feature
     348 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
     350 <400> SEQUENCE: 17
     351 gggggggatcc ggtaccaggt gacgtcgttg ttcagccag                         39
     354 <210> SEQ ID NO: 18
     355 <211> LENGTH: 39
     356 <212> TYPE: DNA
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     360 <220> FEATURE:
     361 <221> NAME/KEY: misc_feature
     362 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
     364 <400> SEQUENCE: 18
     365 gggggggtacc ggatcctcgt agtcggccgc catgacaac                         39
     368 <210> SEQ ID NO: 19
     369 <211> LENGTH: 31
     370 <212> TYPE: DNA
C--> 371 <213> ORGANISM: Artificial/Unknown
     374 <220> FEATURE:
     375 <221> NAME/KEY: misc_feature
     376 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
     378 <400> SEQUENCE: 19
     379 gggggggatcc caggtgacgt cgttgttcag c                                 31
     382 <210> SEQ ID NO: 20
     383 <211> LENGTH: 31
     384 <212> TYPE: DNA
C--> 385 <213> ORGANISM: Artificial/Unknown
     388 <220> FEATURE:
     389 <221> NAME/KEY: misc_feature
     390 <223> OTHER INFORMATION: Description of Artificial Sequence: synthetic DNA
     392 <400> SEQUENCE: 20
     393 gggggggtacc acggtgacgt cgttgttcag c                                 31
     396 <210> SEQ ID NO: 21
     397 <211> LENGTH: 32
     398 <212> TYPE: DNA
C--> 399 <213> ORGANISM: Artificial/Unknown
     402 <220> FEATURE:
     403 <221> NAME/KEY: misc_feature
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Please correct this in subsequent sequences

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 11/23/2004
PATENT APPLICATION: US/09/462,480A TIME: 14:08:32

Input Set : A:\066001650XPCT.txt
Output Set: N:\CRF4\11222004\I462480A.raw

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:34; Xaa Pos. 11

VERIFICATION SUMMARY DATE: 11/23/2004
PATENT APPLICATION: **US/09/462,480A** TIME: 14:08:32

Input Set : **A:\066001650XPCT.txt**
Output Set: **N:\CRF4\11222004\I462480A.raw**

L:301 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:14
L:315 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:15
L:329 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:16
L:343 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:17
L:357 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:18
L:371 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:19
L:385 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:20
L:399 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:21
L:413 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:22
L:427 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:23
L:441 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:24
L:455 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:25
L:469 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
L:483 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:27
L:532 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:29
L:548 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:30
L:564 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:31
L:578 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:32
L:592 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:33
L:620 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34 after pos.:0